```
<110> LAL, Preeti
      AZIMZAI, Yalda
      TANG, Y. Tom
<120 > CYTOCHROME P450 VARIANT
<130> PF-9802 US
<140> To Be Assigned
<141> Herewith
<150>60/218,934
<151> 2000-07-14
<160> 2
<170> PERL Program
<310 > 1
<211> 504
<312> PRT
<213> Homo sapiens
<2220>
<221> misc_feature
<223 > Incyte ID No: 2515666CD1
-:400 > 1
Met Ala Leu Ser Gln Ser Val Pro Phe Ser Ala Thr Glu Leu Leu
                                      1.0
Leu Ala Ser Ala Ile Phe Cys Leu Val Phe Trp Val Leu Lys Gly
                                      25
                 20
Leu Arg Pro Arg Val Pro Lys Gly Leu Lys Ser Pro Pro Gln Pro
                 35
                                      40
Trp Gly Trp Pro Leu Leu Gly His Val Leu Thr Leu Gly Lys Asn
                                      55
                 50
Pro His Leu Ala Leu Ser Arg Met Ser Gln Arg Tyr Gly Asp Val
                 65
                                      70
Leu Gln Ile Arg Ile Gly Ser Thr Pro Val Leu Val Leu Ser Arg
                                      85
                 8.0
Leu Asp Thr Ile Arg Gln Ala Leu Val Arg Gln Gly Asp Asp Phe
                 95
                                     100
Lys Gly Arg Pro Asp Leu Tyr Thr Ser Thr Leu Ile Thr Asp Gly
                1.10
                                     115
Gln Ser Leu Thr Phe Ser Thr Asp Ser Gly Pro Val Trp Ala Ala
                125
                                     130
Arg Arg Arg Leu Ala Gln Asn Ala Leu Asn Thr Phe Ser Ile Ala
                                     1.5
                140
Ser Asp Pro Ala Ser Ser Ser Cys Tyr Leu Glu Glu His Val
```

155

170

185

200

215

230

160

175

190

205

220

235

Ser Lys Glu Ala Lys Ala Leu Ile Ser Arg Leu Gln Glu Leu Met

Ala Gly Pro Gly His Phe Asp Pro Tyr Asn Gln Val Val Val Ser

Val Ala Asn Val Ile Gly Ala Met Cys Phe Gly Gln His Phe Pro

Glu Ser Ser Asp Glu Met Leu Ser Leu Val Lys Asn Thr His Glu

Phe Val Glu Thr Ala Ser Ser Gly Asn Pro Leu Asp Phe Phe Pro

Ile Leu Arg Tyr Leu Pro Asn Pro Ala Leu Gln Arg Phe Lys Ala

15

3.0

75

105

1.0

135

150

```
245
                                      350
Phe Asn Gln Arg Phe Leu Trp Phe Leu Gln Lys Thr Val Gln Glu
                 260
                                      J65
                                                           270
His Tyr Gln Asp Phe Asp Lys His Ser Lys Lys Gly Pro Arg Ala
                275
                                      280
Ser Gly Asn Leu Ile Pro Gln Glu Lys Ile Val Asn Leu Val Asn
                290
                                      295
                                                           300
Asp Ile Phe Gly Ala Gly Phe Asp Thr Val Thr Thr Ala Ile Ser
                                                           315
                                      3.10
                305
Trp Ser Leu Met Tyr Leu Val Thr Lys Pro Glu Ile Gln Arg
                                                          Lys
                320
                                      325
                                                           330
Ile Gln Lys Glu Leu Asp Thr Val Ile Gly Arg Glu Arg Arg Pro
                                     340
                                                           345
                335
Arg Leu Ser Asp Arg Pro Gln Leu Pro Tyr Leu Glu Ala Phe Ile
                                      355
                                                           360
                350
Leu Glu Thr Phe Arg His Ser Ser Phe Leu Pro Phe Thr Ile Pro
                                     370
                                                           375
                365
His Ser Thr Thr Arg Asp Thr Thr Leu Asn Gly Phe Tyr Ile Pro
                380
                                      385
                                                           390
Lys Lys Cys Cys Val Phe Val Asn Gln Trp Gln Val Asn His Asp
                                                           405
                395
                                      400
Pro Glu Leu Trp Glu Asp Pro Ser Glu Phe Arg Pro Glu Arg Phe
                410
                                      415
Leu Thr Ala Asp Gly Thr Ala Ile Asn Lys Pro Leu Ser Glu Lys
                                                           435
                425
                                     430
Met Met Leu Phe Gly Met Gly Lys Arg Arg Cys Ile Gly Glu Val
                440
                                      445
                                                           450
Leu Ala Lys Trp Glu Ile Phe Leu Phe Leu Ala Ile Leu Leu Gln
                                      450
                455
Gln Leu Glu Phe Ser Val Pro Pro Gly Val Lys Val Asp Leu Thr
                                      475
                470
Pro Ile Tyr Gly Leu Thr Met Lys His Ala Arg Cys Glu His Val
                                                           495
                485
                                      490
Gln Ala Arg Leu Arg Phe Ser Ile Asn
                500
```

```
<210> 2
<311> 1790
<312> DNA
<313> Homo sapiens
<320>
```

<221> misc_feature

<223> Incyte ID No: 2515666CB1

<400> 2

```
cagccattac aaccetgeca ateteaagea ectgeeteta cagttggtac agatggeatt 60 gteecagtet gtteeettet eggecacaga getteteetg geetetgeca tettetgeet 120 ggtattetgg gtgeteaagg gtttgaggee tegggteee aaaggeetga aaagteeaee 180 acagecatgg ggetggeet tgetegggea tgtgetgaee etggggaaga accegeaeet 240 ggeaetgtea aggatgagee agegetaegg ggaegteetg cagateegea ttggeteeae 300 geeegtgetg gtgetgagee geetggaeae eateeggeag geeetggtge ggeagggega 360 egattteaag ggeeggeetg acetetacae etceaeeete ateaetgatg geeagagett 420 gaeetteage acagaetetg gaeeggtgt ggetgeeege eggegeetgg eecagaatge 480 eetcaaeaee tteteeateg eetetgaeee agetteetea teeteetget aeetggagga 540 geatgtgage aaggaggeta aggeeetgat eageaggttg eaggagetga tggeagggee 600 tgggeaette gaeeettaea ateaggtggt ggtgteagtg geeaaegtea ttggtgeeat 650 gtgettegga eageaettee etgagagtag egatgagatg eteageeteg tgaagaaeae 720 teatgagtte gtggagaetg eeteeteegg gaaeeeeetg gaeeteetee eeateetteg 780 etaeetgeet aaeeetgeee tgeagaggtt eaaggeette aaeeagaggt teetgtggtt 840 eetegagaaa acagteeagg ageaetatea ggaeetttgae aageaeggge 540 eeteetgeet aaeeetgee tgeagaggtt eaaggeette aaeeagaggt teetgtggtt 840 eetgeagaaa acagteeagg ageaetatea ggaeetttgae aageaegga 540 eetegagagaaa aeagteeagg ageaetatea ggaeetttgae aaeeagaggt teetgtggtt 840 eetgeagaaaa acagteeagg ageaetatea ggaeetttgae aageaeagga agaagggee 900
```

PF-0802 US

ggcaacctca	tcccacagga	gaagattgtc	aaccttgtca	atgacatctt	960
tttgacacag	tcaccacagc	catctcctgg	agcctcatgt	accttgtgac	1020
atacagagga	agatccagaa	ggagctggac	actgtgattg	gcagggagcg	1080
ctctctgaca	gaccccagct	gccctacttg	gaggccttca	tcctggagac	1140
tcctccttct	tgcccttcac	catcccccac	agcacaacaa	gggacacaac	1200
ttctacatcc	ccaagaaatg	ctgtgtcttc	gtaaaccagt	ggcaggtcaa	1260
gagctgtggg	aggacccctc	tgagttccgg	cctgagcggt	tcctcaccgc	1320
					1380
atcggggaag	tcctggccaa	gtgggagatc	ttcctcttcc	tggccatcct	1440
ctggagttca	gcgtgccgcc	gggcgtgaaa	gtcgacctga	cccccatcta	1500
atgaagcacg	cccgctgtga	acatgtccag	gcgcggctgc	gcttctccat	1560
					1620
					1630
					1740
					1790
	tttgacacag atacagagga ctctctgaca tcctccttct ttctacatcc gagctgtggg gccattaaca atcggggaag ctggagttca atgaagcacg agacaccacc gtttctcttc cccagcattt	tttgacacag tcaccacagc atacagagga agatccagaa ctctctgaca gaccccagct tcctccttct tgcccttcac ttctacatcc ccaagaaatg gagctgtggg aggacccctc gccattaaca agcccttgag atcggggaag tcctggccaa ctgaagcaca gcgtgtca agacaccac attctgagc gtttctctt cccagcatt tgggggca	tttgacacag tcaccacage catetectgg atacagagga agatecagaa ggagetggac etetetgaca gacceaget gecetaettg teeteettet tgecetteae cateeceae ttetacatee ecaagaaatg etgtgtette gagetgtggg aggaceette tgagtteegg ecattaaca agecettgag tgagaagatg ateggggaag teetggecaa gtgggggtgaaa atgaagcacg eeegetgtga acatgteeag agacaccace attetettt ttaaaaaata eceageatt tgggaggea ggttgaggg	tttgacacag tcaccacagc catetectgg agcetcatgt atacagagga agatecagaa ggagetggac actgtgattg etetetgaca gaccecaget gecetacttg gaggeettea tectecttet tgecetteac catececac agcacaacaa ttetacatec ecaagaaatg etgtgtette gtaaaccagt gagetgtggg aggacecete tgagtteegg ectgageggt gecattaaca agecettgag tgagaagatg atgetgtttg ateggggaag teetggecaa gtgggagate tteetetee etgagattca gegtgeege gggegtgaaa gtegacetga agacaccace attetgagge cagggagega gtgggggeca gtteetete ettettet ttaaaaaaata geagetttag	atacagagga agatccagaa ggagctggac actgtgattg gcagggagcg ctctctgaca gacccagct gccctacttg gaggcttca tcctggagac tcctctctt tgcccttcac catccccac agcacaacaa gggacacaac ttctacatcc ccaagaaatg ctgtgtcttc gtaaaccagt ggcaggtcaa gagctgtggg aggacccct tgagttccgg cctgagcggt tcctcaccgc gccattaaca agcccttgag tgagaagatg atgctgtttg gcatgggcaa atcggggaag tcctggccaa gtgggagatc ttcctcttc tggcatcct ctggagttca gcgtgccgc gggcgtgaaa gtcgacctga ccccatcta atgaagcacg cccgctgtga acatgtccag gcgggggca gttctctct cttctttt ttaaaaaata gcagctttag ccaagtgcag cccagcatt tgggaggca aggttggagg atcatttgag cccagggtt